# Introduction to the analysis package.

Yo Sato (Tohoku University) Belle II Starter Kit workshop, KEK, January 2020







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Candidate/particle based analysis

**Physics quantities and the VariableManager** 

How to get data out

Nomenclature

One final thing about Belle vs. Belle II



DESY.

DESY.

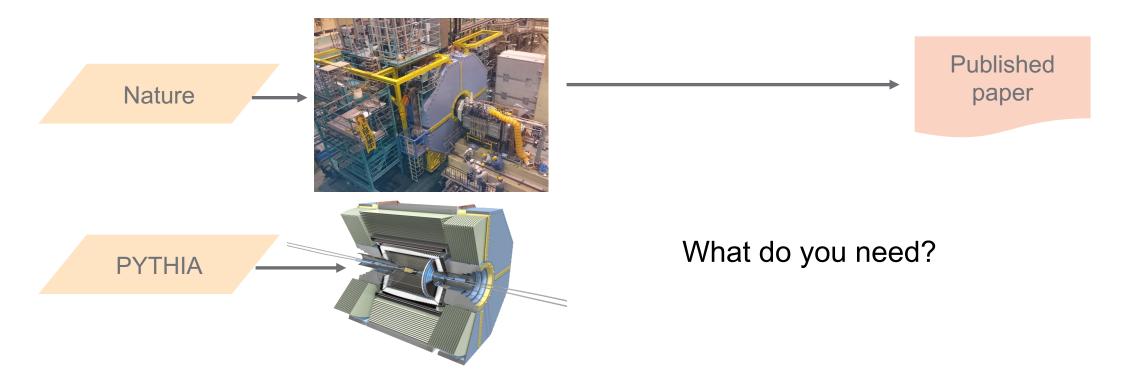
## An experiment's workflow

## The big picture

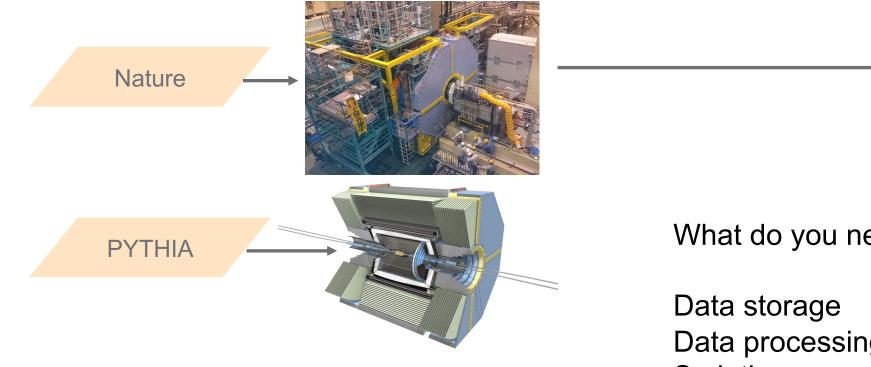
#### Tsukuba Hall









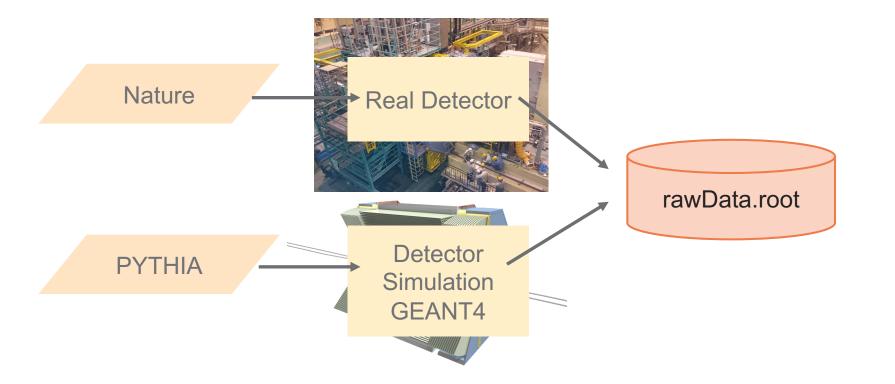


Published paper

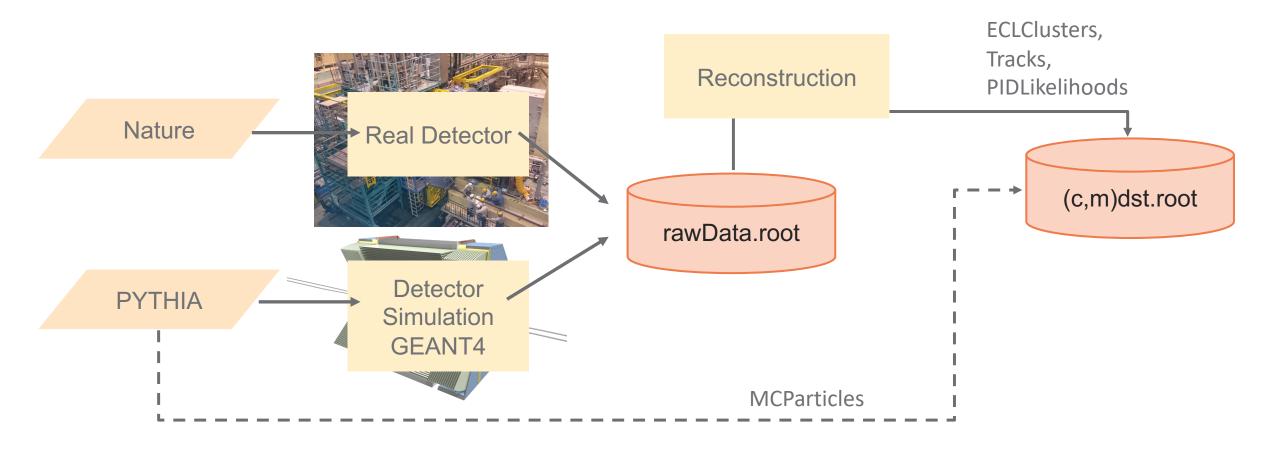
What do you need?

Data storage  $\rightarrow$  root (trees) files Data processing  $\rightarrow$  C++ code Scripting  $\rightarrow$  Python

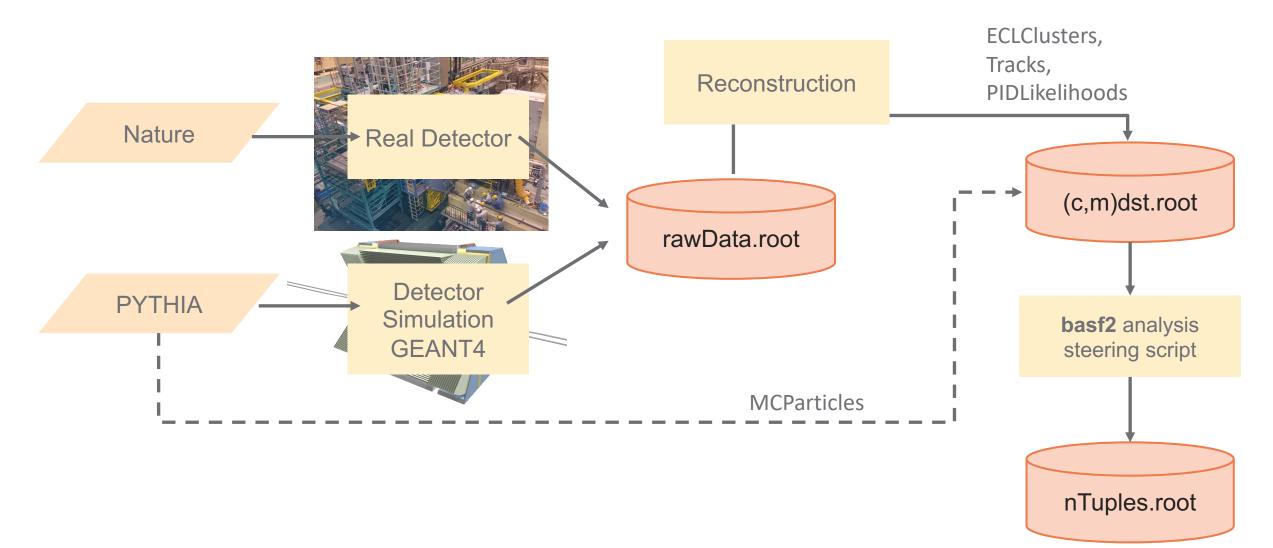




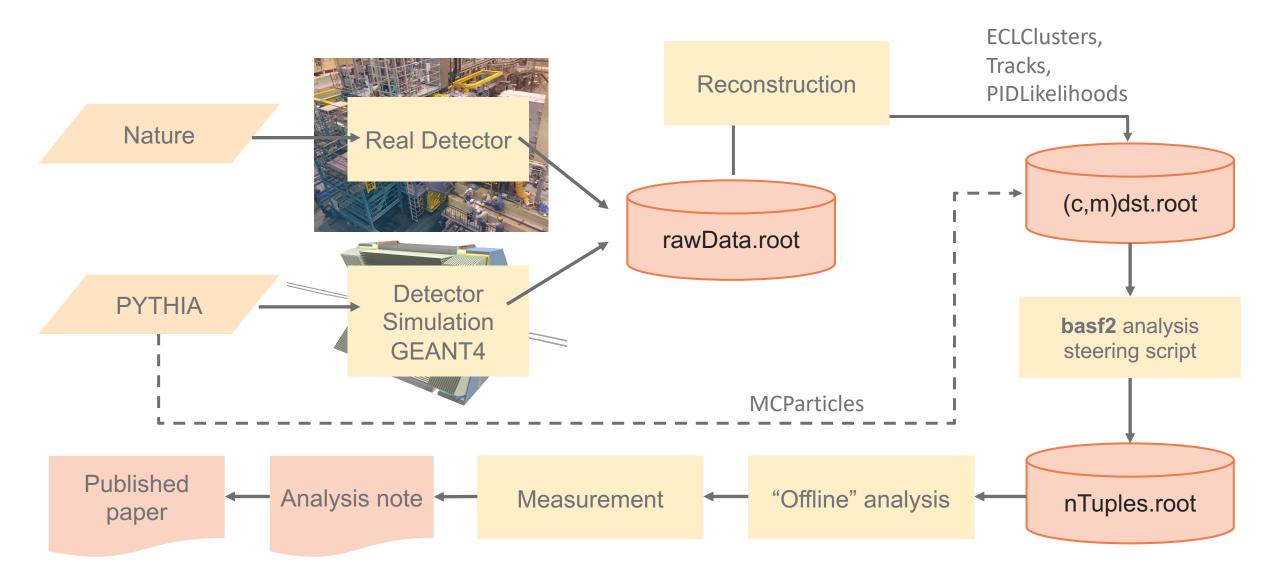




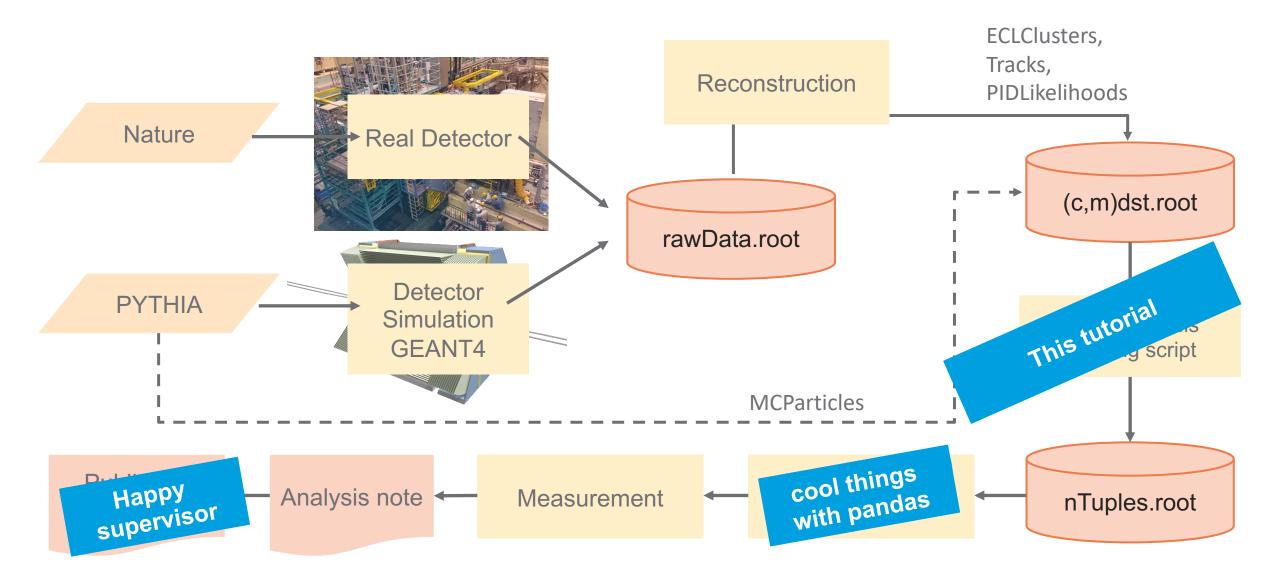












## What is basf2?

DESY.

## The code

basf2 is C++14 "under the hood"

- Packages contain C++ modules to manipulate data.
- In analysis: we have code to build **particles** from primitive objects (like tracks and calorimeter clusters).
- We also calculate physics quantities, and apply cuts.

#### Python 3.6 code for steering

- Load and configure C++ modules
  - analysis modules and modules from other packages
- Also python does *some* high-level analysis tasks.
- You will write a fair bit of python during the workshop.



## The code

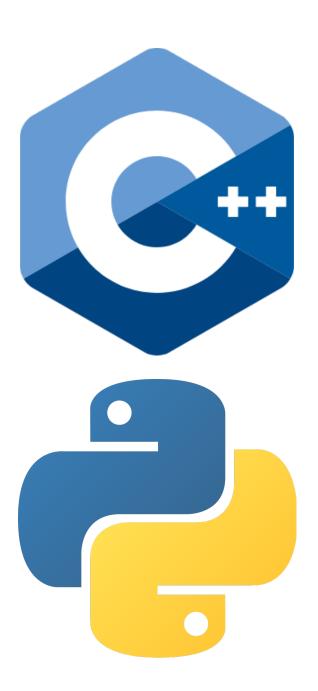
basf2 is C++14 "under the hood"

- In analysis: we have **FFEID** FEED FOR primitive objects (like tracks and called a line of the also calculated and the also cal

#### Python 3.6 code for steering

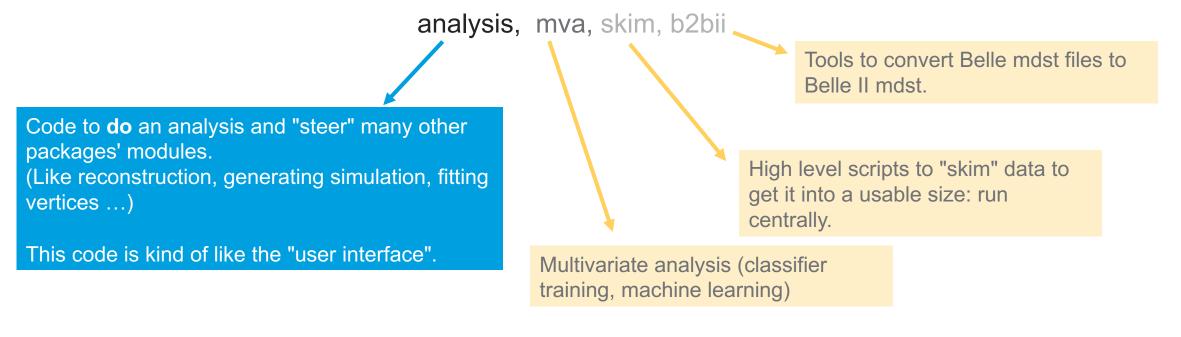
- Load and configure C++ modules
  - analysis modules and m
- Also python does

ves cadable scripts Readable start You will write python during the workshop.



## What is the analysis package?

- Our software is organised into "packages".
  - https://stash.desy.de/projects/B2/repos/software/browse
  - There are packages for subdetectors, tracking, simulation...
  - As a student/postdoc/collaborator you might work on some of them.
- BUT! When you want to do a physics measurement. You really only care about:



## I'm stuck, where do I go for help?

**Probably the most important slide** 

questions for anything really, not just analysis package	https://questions.belle2.org
documentation there is fairly good documentation	https://software.belle2.org
<b>examples</b> so, for us <packagename> = analysis</packagename>	<pre>\$BELLE2_RELEASE_DIR/<packagename>/examples</packagename></pre>
the code	https://stash.desy.de/projects/B2/repos/software/browse \$BELLE2_RELEASE_DIR

## I'm stuck, where do I go for help?

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examples so, for us <packagename> = analysis

the code

https://questions.belle2.org

https://software.belle2.org

Go here first! SE\_DIR/<packagename>/examples

https://stash.desy.de/projects/B2/repos/software/browse **\$BELLE2 RELEASE DIR** 

# Invoking basf2 and the equivalent of "Hello World"

Belle analysis software framework 2

- Let's setup basf2 !
- But, first of all, you should know which release can be and should be used.

```
. /cvmfs/belle.cern.ch/tools/b2setup
 b2setup --help # You can see available releases of basf2
 The following releases are available:
. . .
    release-04-01-00
    release-04-01-01
    light-1912-icarus
 b2help-releases # or you can ask the recommended release
release-04-01-01
```

Belle analysis software framework 2

- Q. Which release should I use?
- A. I really recommend you to use the latest release-AA-BB-CC or light-YYMM-CODENAME.

#### release-AA-BB-CC

A full release, such as release-04-01-01 and release-03-02-04. If you execute `b2help-releases`, it tells the latest full release.

#### light-YYMM-CODENAME

A light release, such as light-1912-icarus and light-1911-heracles.

Made from only a few packages, analysis, mdst, skim, b2bii (from light-1912) etc.. They are suitable for doing analysis!

Belle analysis software framework 2

- Hopefully you've seen this already.
- If not you will see it many times in this workshop.
- At the command line:

```
$ . /cvmfs/belle.cern.ch/tools/b2setup
$ b2setup release-04-01-01
$ basf2 --info
```

Belle analysis software framework 2

- Then Let's setup basf2 for the release!
- `. /cvmfs/belle.cern.ch/tools/b2setup` needs only once per session.

\$ \$ . /cvmfs/belle.cern.ch/tools/b2setup release-04-01-01 basf2 --info \$ \$ \$

## Today we'll use jupyter

#### You will cover command line usage later in the workshop

#### https://jupyterhub.belle2.org

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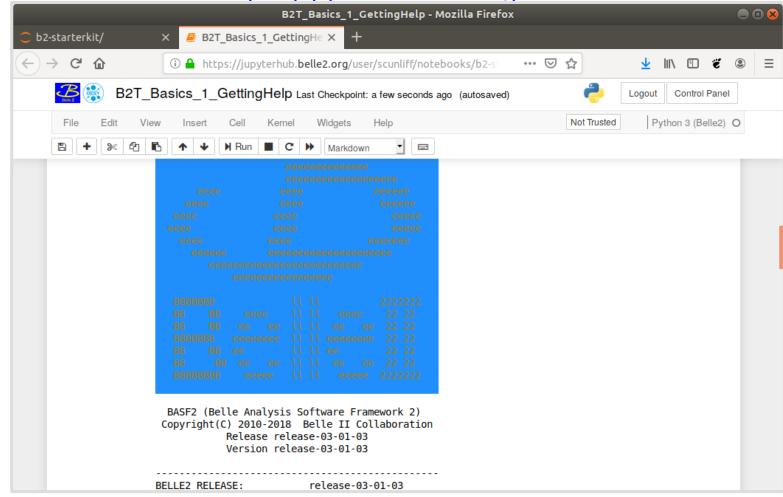
## Start a server, open the first exercise...

#### https://jupyterhub.belle2.org

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Actually if you already know what release you want then you can do it all in once step:			
<pre>source /cvmfs/belle.cern.ch/tools/b2setup release-03-01-01</pre>			
To list all available releases:			
b2setuphelp			
You <b>don't</b> need to do that in this ipython/jupyter environment (we've set it all up for you). But please remen need it later.	mber this, yo	u'll	
In []: !basf2info			

### basf2 --info

#### https://jupyterhub.belle2.org



## basf2 --info

#### Also works on a terminal to KEKCC or NAF

## Source the logon script, pick a release

#### **Execute this**

#### Terminal

#### File Edit View Search Terminal Help

➤ source /cvmfs/belle.cern.ch/tools/b2setup release-03-01-03 Belle II software tools set up at: /cvmfs/belle.cern.ch/tools Environment setup for release: release-03-01-03 Central release directory : /cvmfs/belle.cern.ch/ubuntu1804/releases/release-03-01-03

#### ▶basf2 --info

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### basf2 --info

#### Also works on a terminal to KEKCC or NAF

#### **Useful information**

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## Modules, paths, the DataStore and how to steer them all

What do we need to process the data?

Kuhr, Pulvermacher, Ritter, Hauth, Braun Comput. Softw. Big Sci. 3 (2019) no.1

1) A set of classes (modules) that process the data

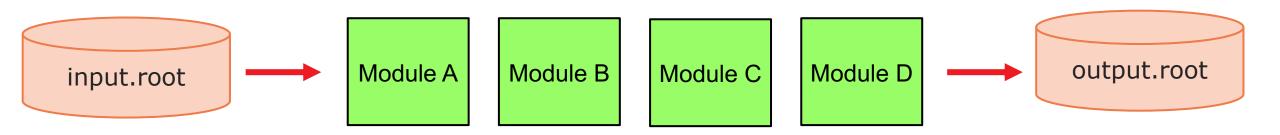
 $\rightarrow \textbf{BASF2 module}$ 

A module is written in C++ or Python and derived from a Module base class that defines the following interface methods:

- initialize(): called before the processing of events to initialize the module.
- beginRun(): called each time before a sequence of events of a new run is processed, e.g., to initialize run-dependent data structures like monitoring histograms.
- event(): called for each processed event.
- endRun(): called each time after a sequence of events of the same run is processed, e.g., to collect runsummary information.
- terminate(): called after the processing of all events.

What do we need to process the data?

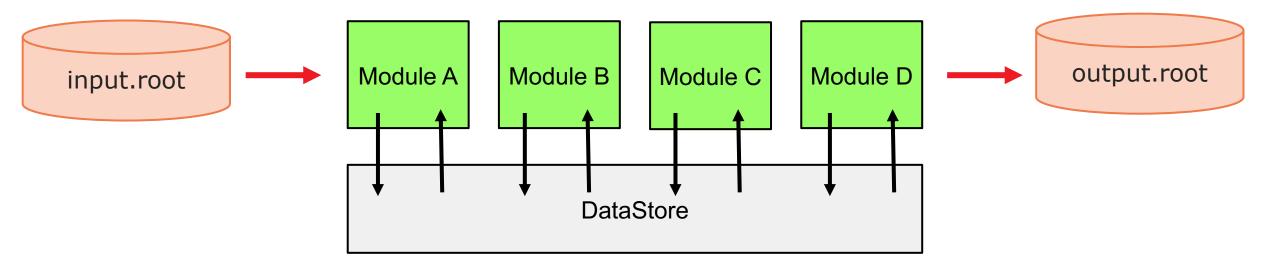
1) A set of classes (modules) that process the data  $\rightarrow$  **BASF2 module** 



What do we need to process the data?

1) A set of classes (modules) that process the data  $\rightarrow$  **BASF2 module** 

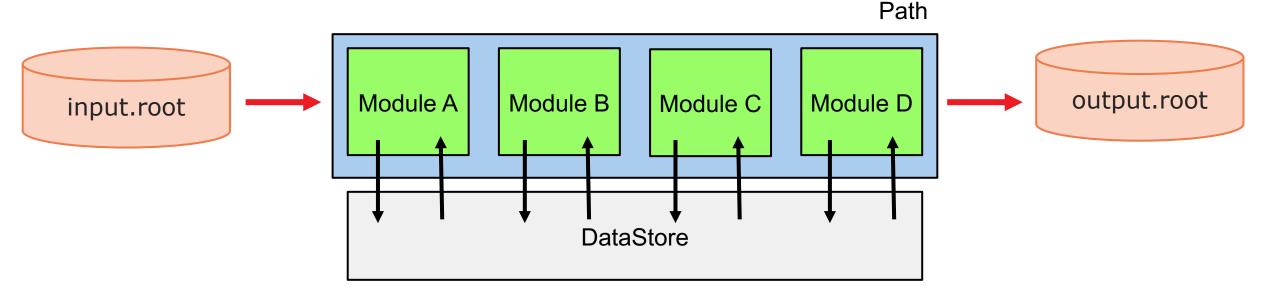
2) A set of classes (dataobjects) that hold the data and allow module to pass thing one to the other  $\rightarrow$  BASF2 dataStore



What do we need to process the data?

1) A set of classes (modules) that process the data  $\rightarrow$  **BASF2 module** 

2) A set of classes (dataobjects) that hold the data and allow module to pass thing one to the other  $\rightarrow$  BASF2 dataStore



3) An order in which the modules must be executed  $\rightarrow$  **BASF2 path** 

## What does a steering file look like?

How to I implement all this?

A steering file is a python script that runs

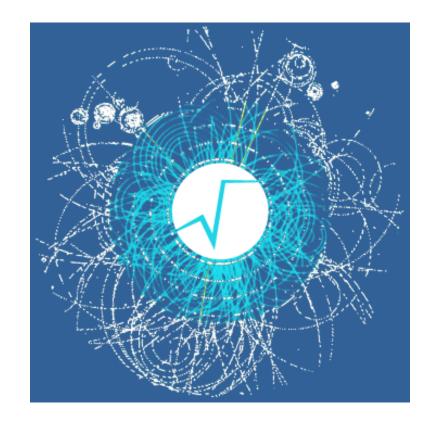
- $\rightarrow$  the modules that you need
- $\rightarrow$  in the order you need
- $\rightarrow$  on the data you need

A word about file types and why you should use the analysis package

## **File types**

#### That basf2 can read and/or create

- A dst contains basf2 objects which will populate a DataStore.
  - data summary table
  - Basically: a special ROOT file.
- The data for physics analysis are "mdst"
  - mini data summary table.
  - Same structure of a dst, but with much less information
  - Input to your analysis package scripts
- The calibration & performance are "cdst"
  - calibration data summary table.
  - mdst + digits
- At the end of your analysis chain you will write out a "normal" root file containing a TTree, TNtuple, or histograms



A relevant question <u>https://questions.belle2.org/question/219</u>

Objects allowed in an *m*dst: <u>https://goo.gl/AB15Ud</u>

## Why use the analysis package?

Can I read the mdst by myself?

mdst are basically root trees containing lists of:

- $\rightarrow$  Track
- $\rightarrow$  TrackFitResult
- $\rightarrow V0$
- $\rightarrow$  PIDLikelihood
- $\rightarrow$  ECLCluster
- $\rightarrow$  KLMCluster
- $\rightarrow$  KIId
- $\rightarrow$  TRGSummary
- → SoftwareTriggerResult
- $\rightarrow$  (MCParticle)

The analysis package has modules to convert these Into more friendly quantities like

- $\rightarrow$  Particle
- $\rightarrow$  ParticleList
- $\rightarrow$  EventShapeContainter
- $\rightarrow$  TagVertex
- $\rightarrow \dots$

 $<sup>\</sup>rightarrow \dots$ 

Can I read the mdst by myself?

Can I open the mdst with my own, custom-made macro and run the analysis?

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Can I open the mdst with my own, custom-made macro and run the analysis?

NO

The mdst contains also the relations bewteen the objects stored in it, which are not trivially handeled by a standalone root macro. **Use always basf2-based code.** 

Can I read the mdst by myself?

N()

Can I open the mdst with my own, custom-made macro and run the analysis?

The mdst contains also the relations bewteen the objects stored in it, which are not trivially handeled by a standalone root macro. **Use always basf2-based code.** 

Should I write my own module that loops over reconstructed objects like the ECLClusters and do the analysis (i.e. Belle-style)?

Can I read the mdst by myself?

Can I open the mdst with my own, custom-made macro and run the analysis?

The mdst contains also the relations bewteen the objects stored in it, which are not trivially handeled by a standalone root macro. **Use always basf2-based code.** 

Should I write my own module that loops over reconstructed objects like the ECLClusters and do the analysis (i.e. Belle-style)?

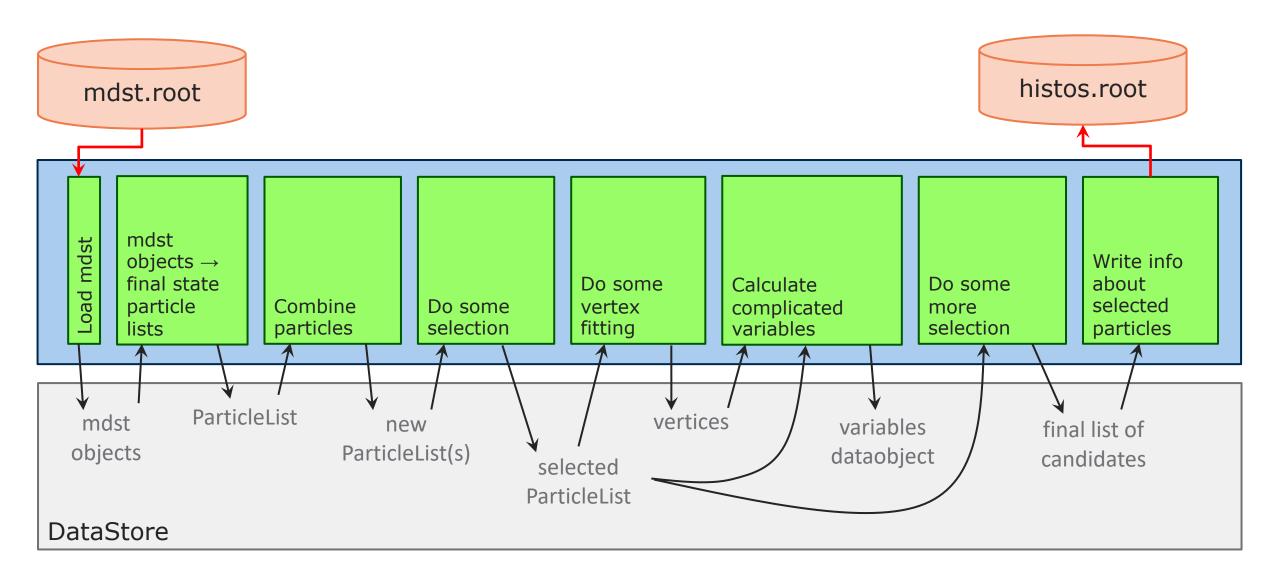
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The relation bewteen analysis object (particles) and the reconstructed objects is not always trivial. One particle may have many trackFitResults The ECLClusters are not photons. Use the modules provided by a detector expert

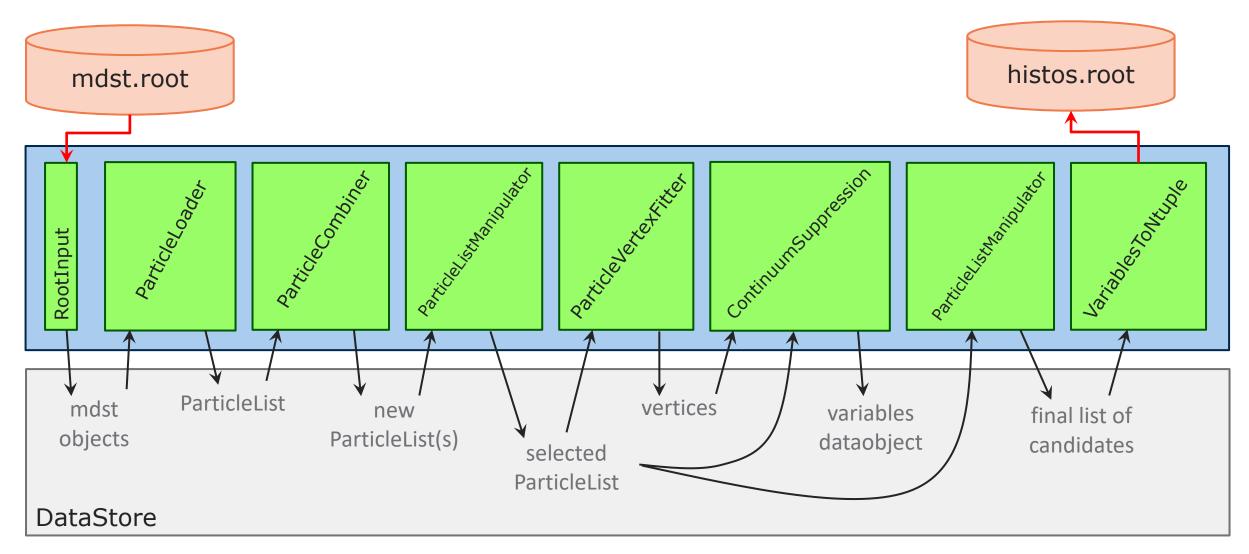
# Now let's step through an example

### A typical path for an analysis job



# A typical path for an analysis job

#### Now with the real names for the modules



#### How to make a path and load a module

• First of all, you have make your path.

```
import basf2
```

```
mypath = basf2.Path()
# mypath basf2.create_path() # Both are OK!
```

#### How to make a path and load a module

- Call a function to load a module and add it to your path.
- That is all you need to do!

```
from modularAnalysis import fillParticleList
```

```
fillParticleList('pi+:highMom', 'p > 1', path=mypath)
```

Load a module to fill a list of particle, ParticleList. We'll see details of ParticleList later.

• What is happening in the function?

#### **The ParticleLoader**

An analysis module which loads particles

fillParticleList('pi+:highMom', 'p > 1', path=mypath)

```
pload = register_module('ParticleLoader')
pload.param('decayStringsWithCuts', [('pi+:highMom', 'p > 1')])
mypath.add_module(pload)
```

- The function wraps these steps in a line.
- Of course, you can write them by hand.
  - But quite quickly scripts become unreadable

#### **The ParticleLoader**

An analysis module which loads particles

- The procedure is same no matter what module you want in path.
  - "Register" the module you want
  - Set "param"eters to the module
  - "Add" the module to you path

```
pload = register_module('ParticleLoader')
pload.param('decayStringsWithCuts', [('pi+:highMom', 'p > 1')])
mypath.add_module(pload)
```

# How to find module documentation

In general

• At the command line:

\$ basf2 -m
\$ basf2 -m ParticleLoader

Take a look at B2T\_Basics\_1\_GettingHelp.ipynb on jupyterhub

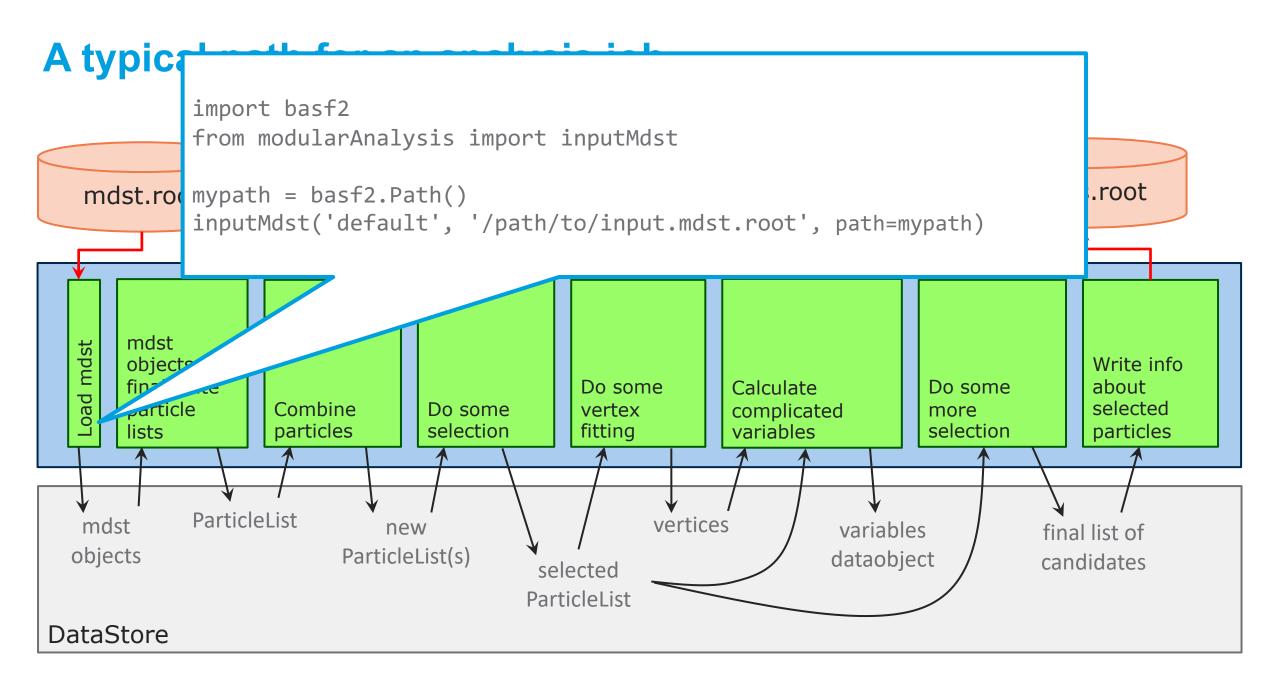
Q: Can you find the **source code** for ParticleLoader? What do you notice about the C++ class name? How does this compare to the module name?

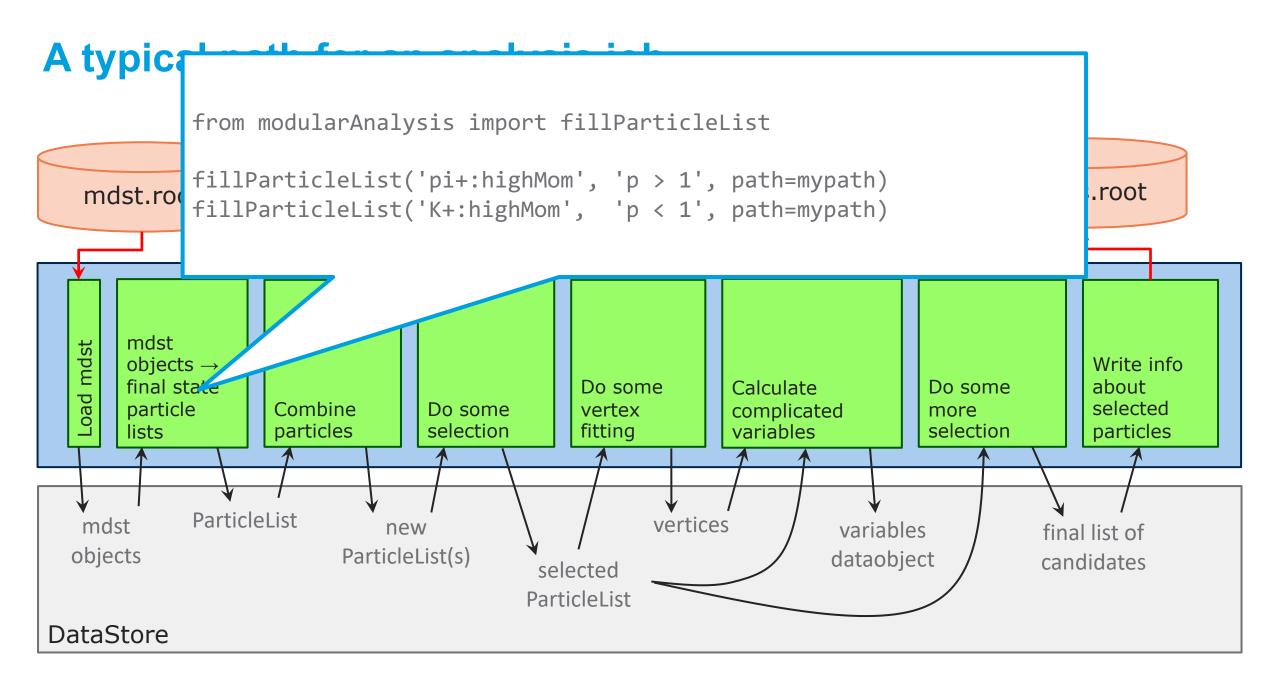
#### **Wrapper functions**

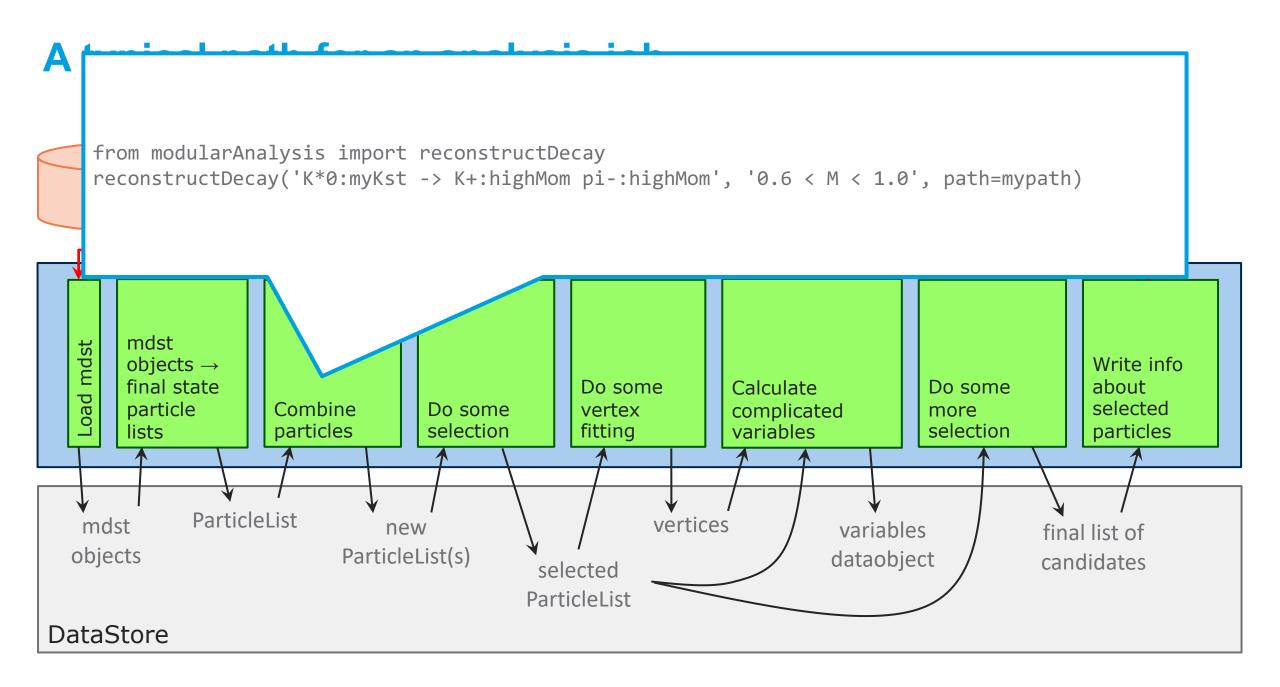
- Functions such as fillParticleList are found in <package>/scripts/\*.py
- The script that you will probably use a lot is analysis/scripts/modularAnalysis.py

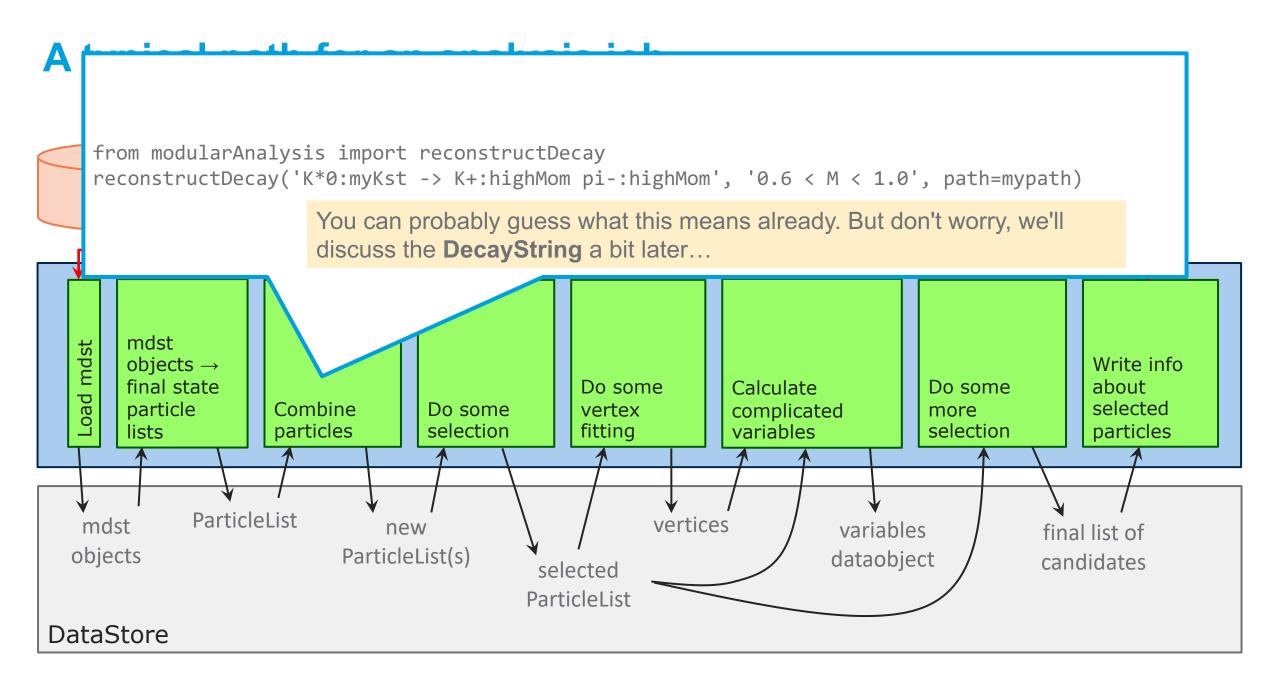
```
# analysis/scripts/modularAnalysis.py
from modularAnalysis import fillParticleList
fillParticleList('pi+:highMom', 'p > 1', path=mypath)
```

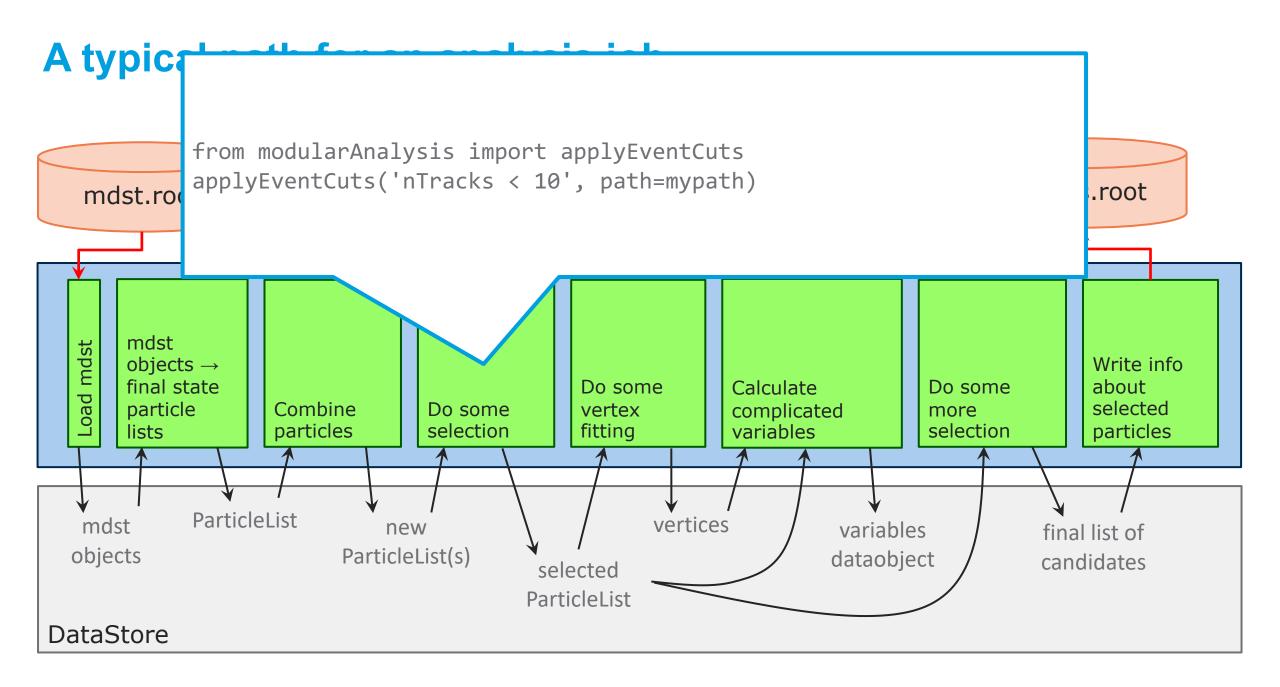
```
# analysis/scripts/vertex.py
from vertex import fitVertex
fitVertex('K*0:myKst', conf_level=0.0, path=mypath)
```

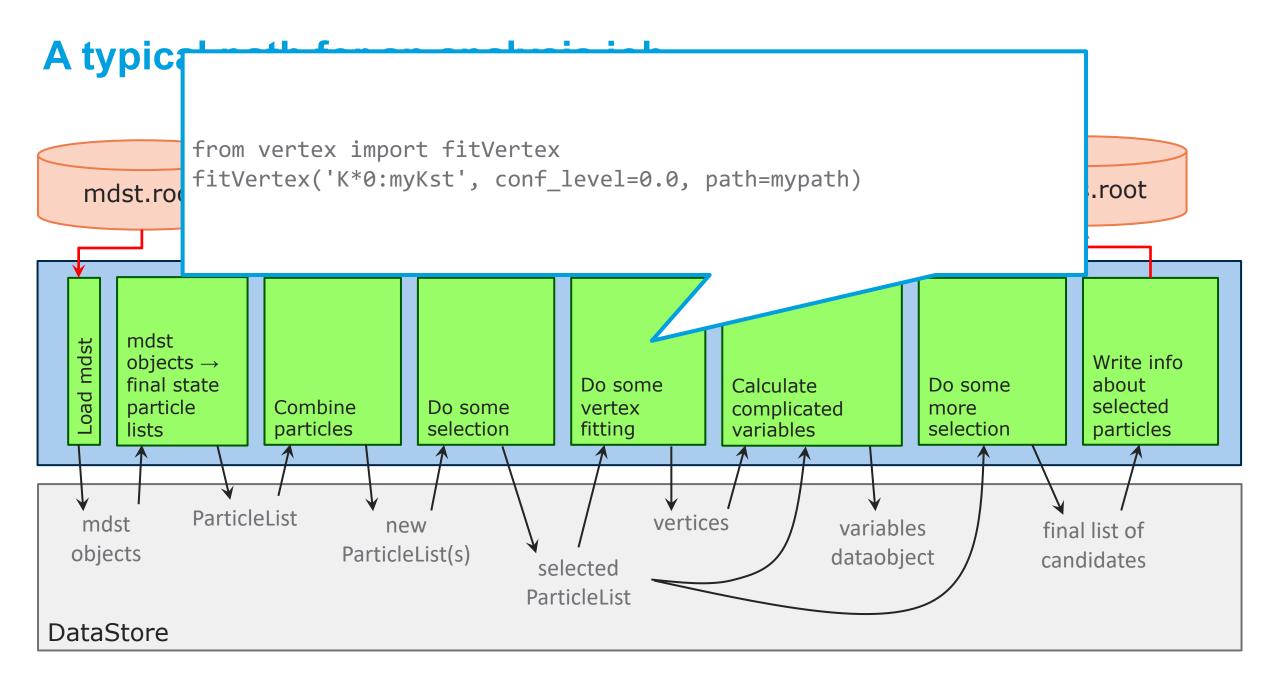




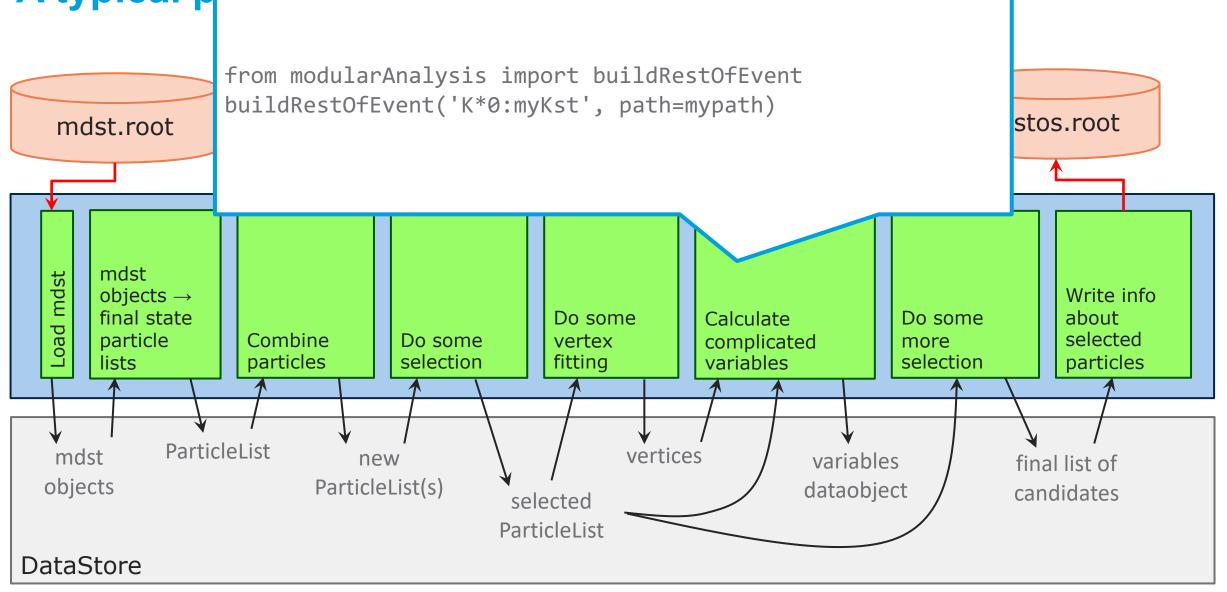




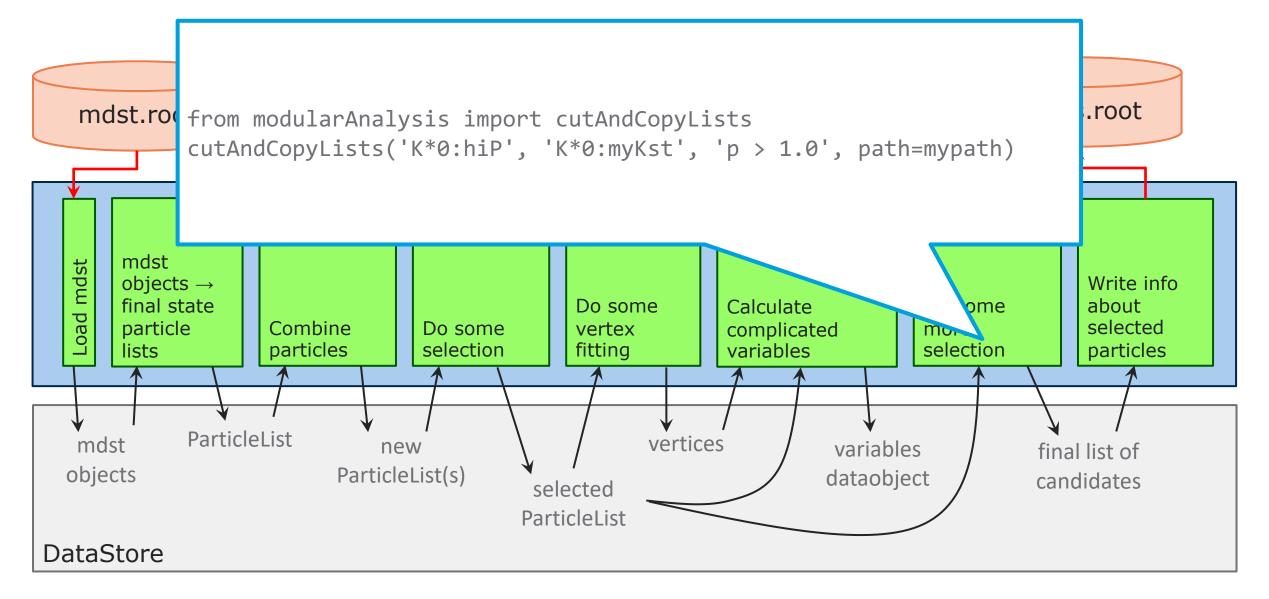


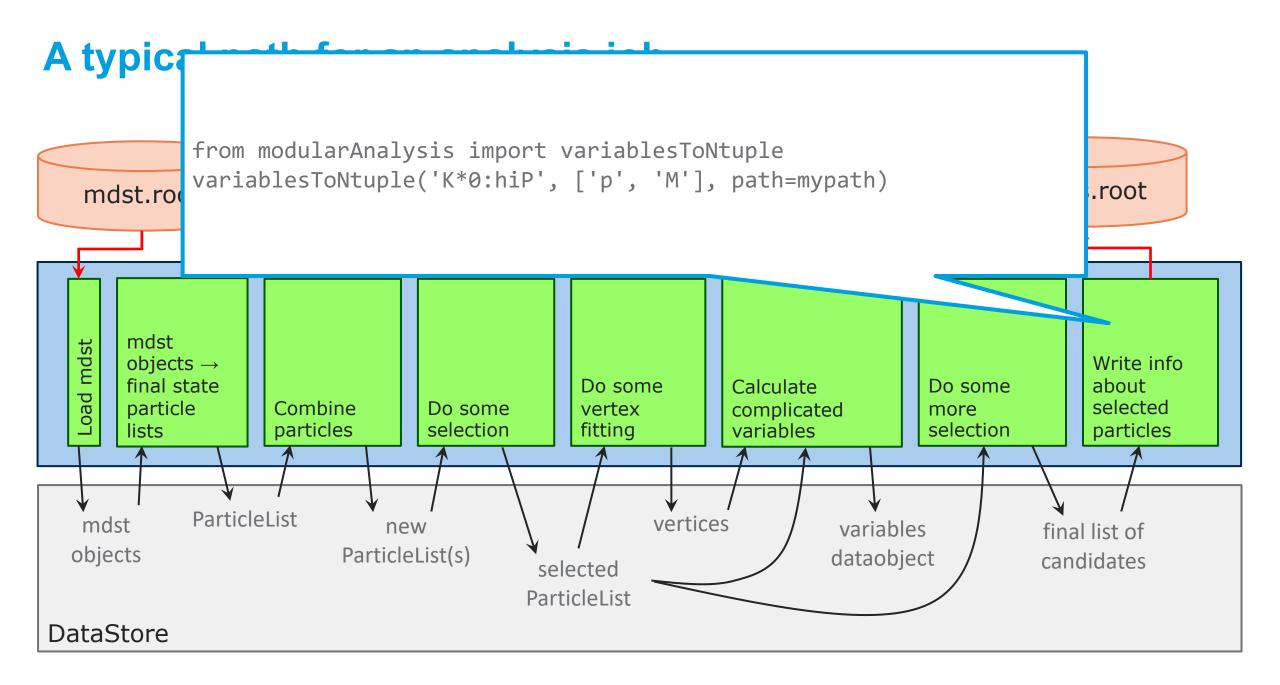


#### A typical path for an analysis job



#### A typical path for an analysis job





#### A template steering script

```
#!/usr/bin/env python3
# -*- coding: utf-8 -
```

import basf2
import modularAnalysis as ma

```
# create a path
mypath = basf2.Path()
```

```
# input mdst file
ma.inputMdst('default','inputMdstFile.root', path=mypath)
```

# PUT YOUR CODE HERE

# process the events
basf2.process(mypath)
# print out the summary
print(basf2.statistics)

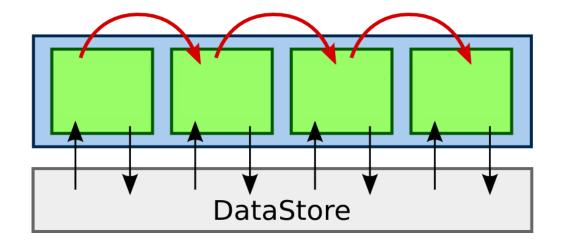


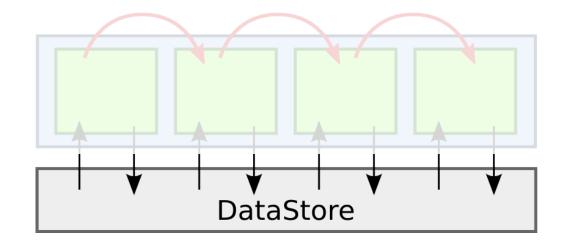
### A template steering script

```
#!/usr/bin/env python3
# -*- coding: utf-8 -
import basf2
import modularAnalysis as ma
# create a path
mypath = basf2.Path()
                                                              Q: what does 'default' mean?
# input mdst file
                                                              What is the argument called?
ma.inputMdst('default','inputMdstFile.root', path=mypath)
                                                              Can you find the documentation?
                                                              Is there a questions post?
# PUT YOUR CODE HERE
# process the events
basf2.process(mypath)
# print out the summary
print(basf2.statistics)
```

# Candidate/particle based analysis

#### Before we get started, here's this diagram again





You can take a look at examples in <package>/dataobjects

- The datastore contains the **dataobjects**
- At the level of analysis, the main dataobject is: ParticleList

#### **Particle-based paradigms**

In the wild

# Event based analysis frameworks Babar, ATLAS, ILC

Particle/candidate based analysis frameworks LHCb, Belle II

#### **Particle-based analysis**

- Take particle lists
- Build up decay parents from daughters
- Make *candidate*s for your decay of interest
- Filter/cut/keep.
- You might have more than one candidate per event.
  - We deal with this after the fact.
  - This is fine. I promise.
  - <u>https://arxiv.org/abs/1703.01128</u>

#### **The Particle class**

It's not crucial to understand the details

- A common representation of all particle types
  - Charged:  $e / \mu / \pi / K / p / d$
  - γ
  - K<sup>0</sup><sub>L</sub>, n
  - K<sup>0</sup><sub>S</sub>, Λ<sup>0</sup>, γ
  - Composite particles:  $\pi^0 / K_S^0 / D / B$

[built up from track + hypothesis] [built up from ECLClusters + !Track] [built up from KLMlusters + ECLClusters + !Track ] [built up from V0 (2 tracks)] [built up from combinations]

- Data members of the class are common to all particle types: mass, momentum, position, PDG code, ...
- Information which is only relevant to certain kinds of particle is saved in separate **analysis package dataobjects** and accessible by **relations**.
  - e.g. ContinuumSuppression
  - .... FlavorTaggerInfo

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- Information which is only relevant to certain kinds of particle is saved in separate **analysis package dataobjects** and accessible by **relations**.
  - e.g. ContinuumSuppression
  - .... FlavorTaggerInfo

Q: what is the name of the module to combine particles?

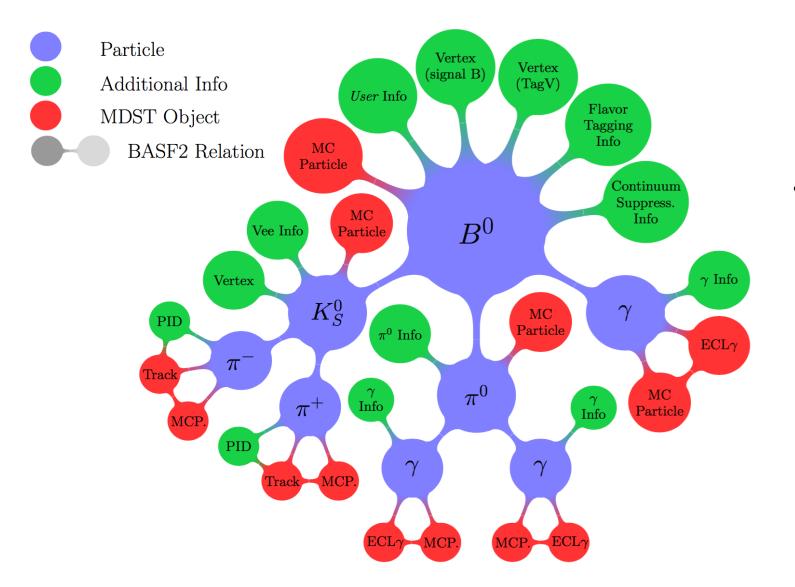
Q: which things on this slide are dataobjects?

#### **ParticleList**

- A group of all particles and anti-particles that belong together logically.
  - e.g.  $K^{*0}$  s (decaying to  $K^{\pm}$  and  $\pi^{\mp}$  with invariant mass in a certain window)
- Can only store particles of the same PDG code (can be different decay modes).
- Doesn't have ownership of the Particle objects.
- ParticleList is the dataobject on which analysis modules operate.

- The physics-performance group provides Standard Particle Lists for which quality benchmarks exist, and systematics will be provided.
  - Currently you may need to optimize selection criteria of ParticleLists by yourself.
  - This will be really recommended to use in future.

#### Some more details on Particles, ParticleLists and Relations



- At each stage we build relations between the dataobjects
  - Like vertex information, ContinuumSuppression
     → all related to Particles
  - Particles themselves related to primitive mdst objects (clusters, tracks)

#### **Fill a ParticleList**

There are two possible ways to fill a list of **stable\* particles** 

1) fill it by hand

import modularAnalysis as ma

ma.fillParticleList('pi+:highMom', 'p > 1', path=mypath)

Q: What is a stable particle in this context?

#### **Fill a ParticleList**

There are two possible ways to fill a list of stable\* particles

1) fill it by hand

import modularAnalysis as ma

ma.fillParticleList( pi+:highMom', 'p > 1', path=mypath)

Particle name. Tells basf2 if the list has to be created from tracks, ELCClusters, KLMClusters or V0 Must given in evt.pdl (or b2help-particles)

evt.pdl = \$BELLE2\_EXTERNALS\_DIR/share/evtgen/evt.pdl

#### **Fill a ParticleList**

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Label name.

Used to distinguish different lists of the same particle type: you choose the name. (But maybe call it something helpful)

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Used to distinguish different lists of the same particle type: you choose the name. (But maybe call it something helpful)

ParticleList name Same label name can be set to another particle name. e.g. pi+:highMom, K+:highMom

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Label name.

Used to distinguish different lists of the same particle type: you choose the name. (But maybe call it something helpful)

Selection criteria Cuts used to select this list. **More on this later**  List name Same label name can be set to another particle name. e.g. pi+:highMom, K+:highMom

There are two possible ways to fill a list of stable\* particles

1) fill it by hand

import modularAnalysis as ma

```
ma.fillParticleList('pi+:highMom', 'p > 1', path=mypath)
```

Once you call the function, pi-:highMom is created automatically. You **don't** have to call

```
ma.fillParticleList('pi-:highMom', 'p > 1', path=mypath)
```

There are two possible ways to fill a list of stable\* particles

1) fill it by hand

import modularAnalysis as ma

ma.fillParticleList('pi+:highMom', 'p > 1', path=mypath)

#### 2) Use the standard particles lists with pre-defined names and cuts

from stdPhotons import stdPhotons
from stdCharged import stdK

stdPhotons('all', path=mypath) # --> gamma:all
stdK('95eff', path=mypath) # --> K+:95eff

First argument will be a label name of the particle list.

The last of the infrastructure things, I promise

- You've already seen the decay string in action.
- Now we can formally describe what this line is doing:

reconstructDecay('K\*0:myKst -> K+:highMom pi-:highMom', '0.6 < M < 1.0', path=mypath)</pre>

- You've already seen the decay string in action.
- Now we can formally describe what this line is doing:

reconstructDecay(	'K*0:myKst -> K+:highMom pi-:highMom'	'0.6 < M < 1.0', path=mypath)
	Decay string	

- You've already seen the decay string in action.
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reconstructDecay(	K*0:myKst -> K+:highMom pi-:highMom	'0.6 < M < 1.0', path=mypath)
	ParticleList ParticleList ParticleList	
	Decay string	

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reconstructDecay	'K*0:myKst -> K+:highMom pi-:highMom'	'0.6 < M < 1.0', path=mypath)
	Arrow (Indicate K*0 decays to K+ pi-)	
	Decay string	

- You've already seen the decay string in action.
- Now we can formally describe what this line is doing:

reconstructDecay(	K*0:myKst -> K+:highMom pi-:highMom	'0.6 < M < 1.0', path=mypath)
	mother arrow daughter0 daughter1	
	Decay string	

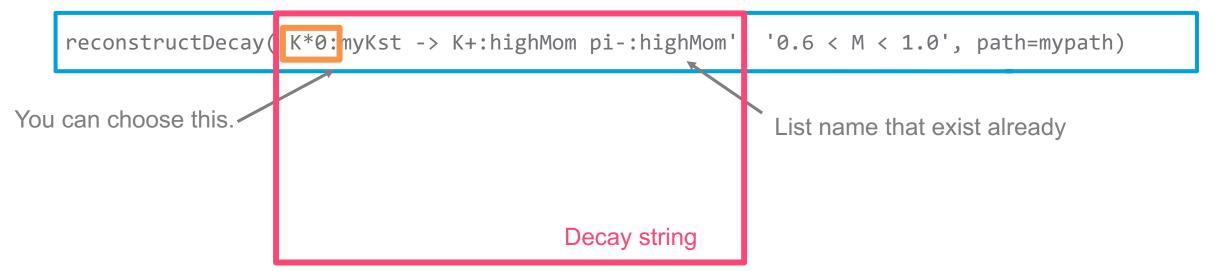
The last of the infrastructure things, I promise

- You've already seen the decay string in action.
- Now we can formally describe what this line is doing:

reconstructDecay	(K*0:myKst -> K+:highMom pi-:highMom'	'0.6 < M < 1.0', path=mypath)
	Particle name. Of your composite particle. Must given in evt.pdl (or b2help-particles) Decay string	

The last of the infrastructure things, I promise

- You've already seen the decay string in action.
- Now we can formally describe what this line is doing:



reconstructDecay() : Reconstruct mother particle from daughters with a given DecayString.
 List names for daughters must exist already. You can choose a list name of a mother.

You don't have to call reconstructDecay('anti-K\*0:myKst -> K-:95eff pi+:95eff', '0.6 < M < 1.0', path=mypath)



One final thing...

- Sometimes you will need to select a particle within a decay string.
- This is done by "carat": ^

'K\*0:myKst -> ^K+:highMom ^pi-:highMom'

• You will see this in action in the exercises later.

# Physics quantities and the Variable Manager

### The VariableManager

It manages variables

- VariableManager is a place in the analysis package to store variables
  - physics quantities: invariant mass, beam-constrained mass, E, p, p<sub>T</sub>, θ, φ, highest energy in a cluster
  - counters: event\_number, nhits, i\_candidate

 Every variable takes at least a Particle\* as input and returns a double (even integer counters like event\_number where this doesn't make much sense)



### The VariableManager

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- VariableManager is a place in the analysis package to store variables
  - physics quantities: invariant mass, beam-constrained mass, E, p, p<sub>T</sub>, θ, φ, highest energy in a cluster
  - counters: event\_number, nhits, i\_candidate

 Every variable takes at least a Particle\* as input and returns a double (even integer counters like event\_number where this doesn't make much sense)

- You've already seen it in **python**, when we used a cut on "M".
  - This comes from the VariableManager.

reconstructDecay('K\*0:myKst -> K+:highMom pi-:highMom', '0.6 < M < 1.0', path=mypath)</pre>



## Where can I get the documentation?

• At the command line:

• Online, the documentation is <u>software.belle2.org</u>.

Take a look at B2T\_Basics\_1\_GettingHelp.ipynb on jupyterhub

Q: What is the beam-constrained mas,  $M_{bc}$  called in the VariableManager?

#### Have you been listening?

Q: Can you figure out the cut string and code to reconstruct B0 -> K\*0 gamma? Supposing you have already reconstructed K\*0:myKst.

import modularAnalysis as ma

```
ma.fillParticleList('pi+:highMom', 'p > 1', path=mypath)
ma.fillParticleList('K+:highMom', 'p > 1', path=mypath)
ma.reconstructDecay('K*0:myKst -> K+:highMom pi-:highMom', '0.6 < M < 1.0', path=mypath)</pre>
```

Q: How can we fill a particle list of gamma?

Q: How can we reconstruct B0 from K\*0 and gamma?

Q: Can you apply following selection on the B0 candidates?  $M_{bc} > 5.2\,{\rm GeV}/c^2$ 

#### Aliases

#### Are awesome and you should use them

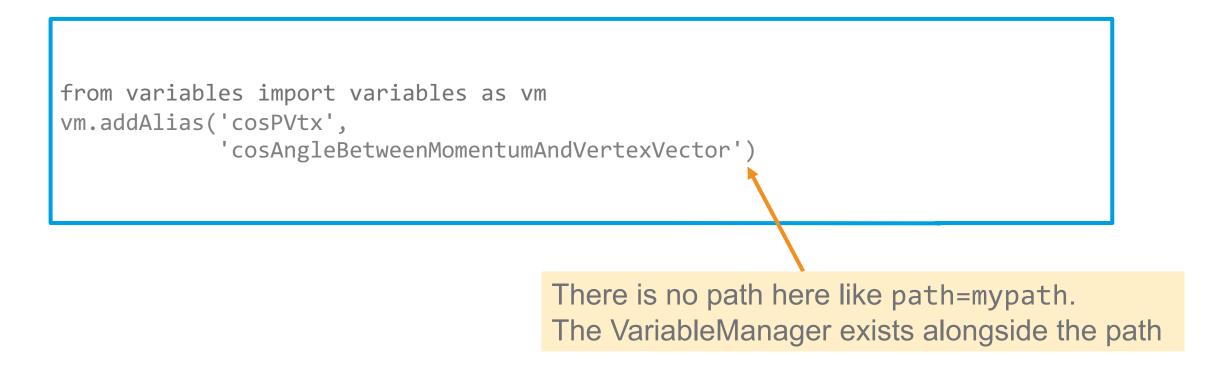
- With more advanced variables, we deliberately give them verbose names in the VariableManager.
- It's important to be clear.
- Some examples:
  - cosAngleBetweenMomentumAndVertexVector
  - totalPhotonEnergyOfEvent
  - trackFindingFailureFlag
- You can also use formula(), abs(), cos() for simple math:
  - formula(missingEnergyOfEventCMS/Ecms)

• ...

#### Aliases

Are awesome and you should use them

- For your code, you can make short aliases.
- This makes your offline data more manageable, and can help with code readability.



# How to get data out

#### Flat ntuples and histograms

**Getting output data** 

There are some modules to store variables.

VariablesTo\*

- a) VariablesToNtuple
- b) VariablesToHistogram
- c) VariablesToEventBasedTree

#### Flat ntuples and histograms

#### **Getting output data**

There are some modules to store variables.

VariablesTo\*

- a) VariablesToNtuple
- b) VariablesToHistogram
- c) VariablesToEventBasedTree

from modularAnalysis import variablesToNtuple

from modularAnalysis import variablesToHistogram

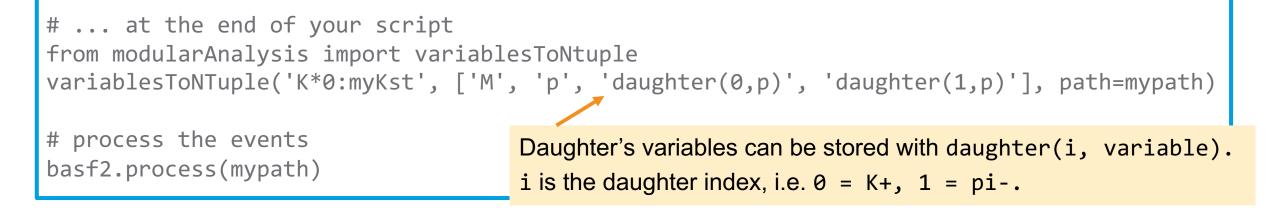
#### **VariablesToNtuple**

- You want to store physics quantities for one ParticleList from the VariableManager.
- Get an ttree (ntuple) of candidates (one row per candidate).

## VariablesToHistogram

- Perhaps you just want a quick histogram.
- Candidate information not preserved.

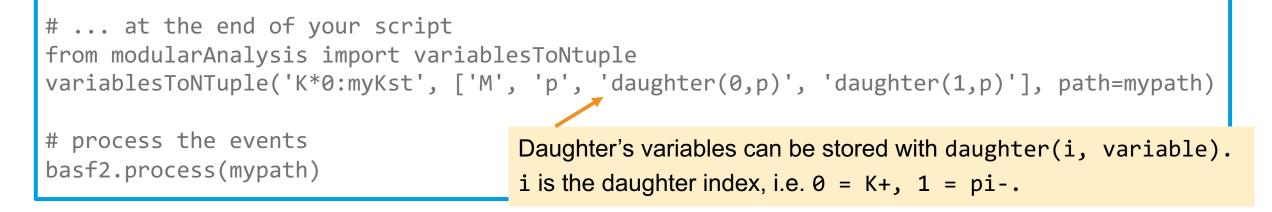
#### **VariablesToNtuple**



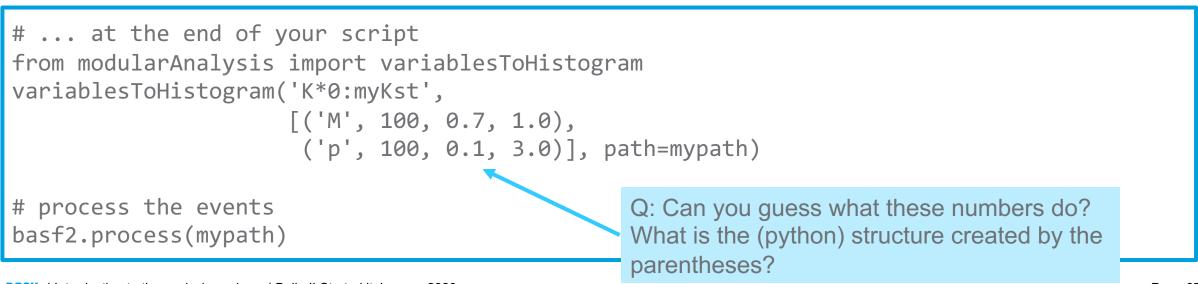
#### VariablesToHistogram

```
# ... at the end of your script
from modularAnalysis import variablesToHistogram
variablesToHistogram('K*0:myKst',
                         [('M', 100, 0.7, 1.0),
                          ('p', 100, 0.1, 3.0)], path=mypath)
# process the events
basf2.process(mypath)
```

#### **VariablesToNtuple**



#### VariablesToHistogram



#### **Utilities**

Helper functions to make daughter's aliases.

- You can store variables of daughters with daughter(i, variable) in a same ntuple.
- You may also want to short aliases for these variables.

```
reconstructDecay('K*0:myKst -> K+:highMom pi-:highMom', '0.6 < M < 1.0', path=mypath)
from variables import variables as vm
vm.addAlias('K_M', 'daughter(0,M)')
vm.addAlias('K_p', 'daughter(0,p)')
vm.addAlias('pi_M', 'daughter(1,M)')
vm.addAlias('pi_p', 'daughter(1,p)') # You may want to add more aliases
vars = ['M', 'p', 'K_M', 'K_p', 'pi_M', 'pi_p']
variablesToNtuple('K*0:myKst', vars, path=mypath)</pre>
```

#### **Utilities**

Helper functions to make daughter's aliases.

• A helper function creates aliases from a list of variables and a DecayString with carat '^'.

All aliases can be checked with printAliases().

# Nomenclature

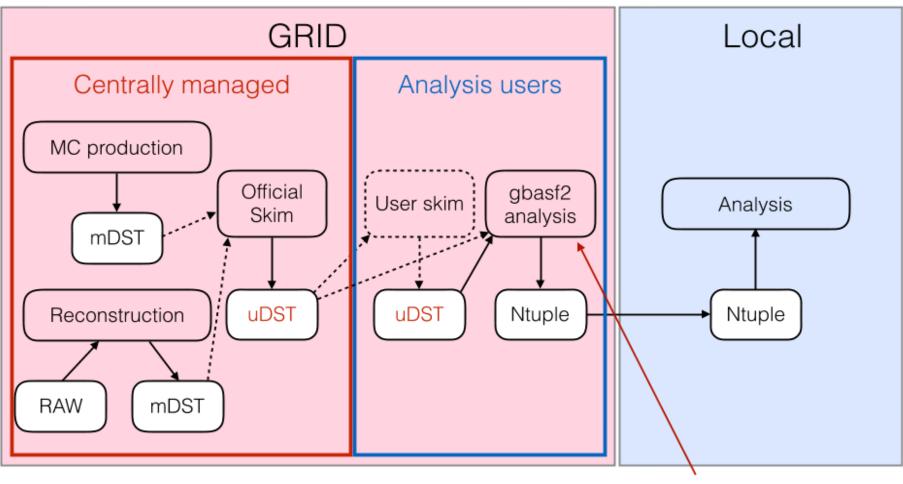
#### Nomenclature

https://confluence.desy.de/ display/BI/Main+Glossary

- **Experiment** (chunk of data-taking ~months).
- Run (chunk of data-taking w/ stable beams ~hours),
- Event.
- **TRG** the hardware trigger (group, device, DAQ)
- L1 the hardware trigger (used interchangeably)
- SoftwareTrigger / **HLT** (the softeare trigger)
- **basf2** "Belle 2 analysis software framework" "the software"
- gbasf2 "The grid job submission tool" "computing"

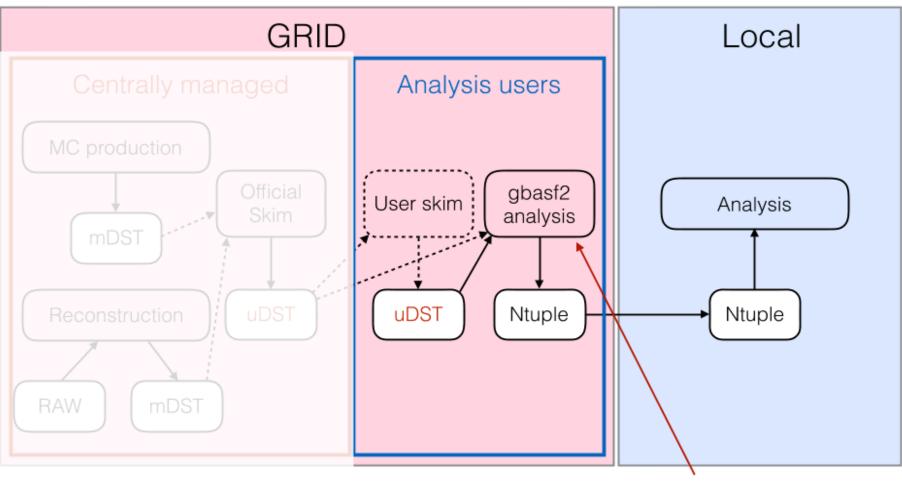
# One final thing about Belle vs. Belle II

#### The Belle II analysis model



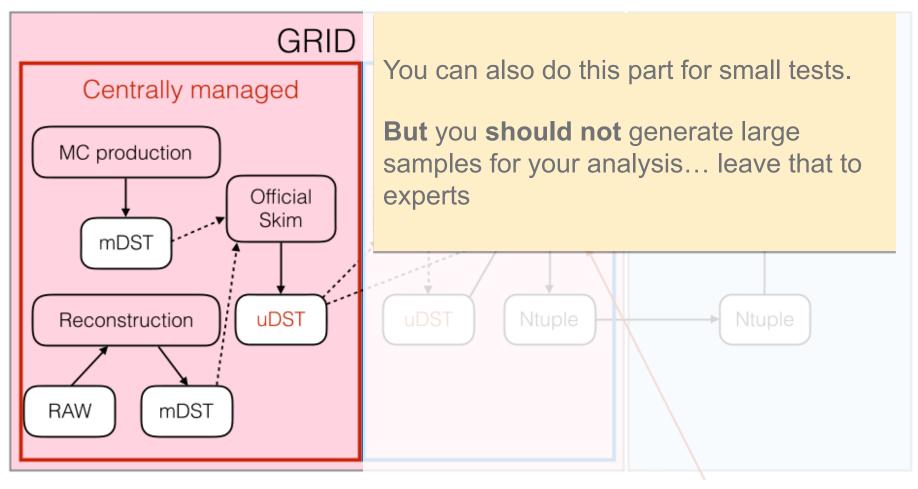
You will learn this tomorrow

#### The Belle II analysis model



You will learn this tomorrow

#### The Belle II analysis model



You will learn this tomorrow

#### Acknowledgements

• Much of the material for these slides (and ideas for how to present it) has been stolen from:

#### Jake Bennett and Anže Zupanc

## **Versions of this tutorial material**

- Version 1: Sam (February 2018).
- Version 2: Ilya and Sam (June 2018).
- Version 3: Sam (November 2018).
- Version 4: Umberto (February 2019)
- Version 5: Hannah and Sam (June 2019)
- Version 6: Yo and Ilya (January 2020)